

# SEQUENCE LISTING

<110> Chen et al.

<120> METHODS AND COMPOSITIONS FOR STIMULATING AXON REGENERATION AND PREVENTING NEURONAL CELL DEGENERATION

<130> ERM-105.01

<160> 4

<170> PatentIn version 3.0

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<212> DNA

<213> homo sapiens

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<221> CDS

<222> (32)..(751)

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Gly Tyr Asp Asn Arg Glu Ile Val Met Lys Tyr Ile His Tyr Lys Leu  
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tcg cag agg ggc tac gag tgg gat gcg gga gat gtg ggc gcc gcg ccc 148  
Ser Gln Arg Gly Tyr Glu Trp Asp Ala Gly Asp Val Gly Ala Ala Pro  
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ccg ggg gcc gcc ccc gcg ccg ggc atc ttc tcc tcg cag ccc ggg cac 196  
Pro Gly Ala Ala Pro Ala Pro Gly Ile Phe Ser Ser Gln Pro Gly His  
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acg ccc cat aca gcc gca tcc cgg gac ccg gtc gcc agg acc tcg ccg 244  
Thr Pro His Thr Ala Ala Ser Arg Asp Pro Val Ala Arg Thr Ser Pro  
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ctg cag acc ccg gct gcc ccc ggc gcc gcc gcg ggg cct gcg ctc agc 292  
Leu Gln Thr Pro Ala Ala Pro Gly Ala Ala Ala Gly Pro Ala Leu Ser  
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Pro Val Pro Pro Val Val His Leu Thr Leu Arg Gln Ala Gly Asp Asp  
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Phe Ser Arg Arg Tyr Arg Arg Asp Phe Ala Glu Met Ser Arg Gln Leu  
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cac ctg acg ccc ttc acc gcg ccg gga cgc ttt gcc acg gtg gtg gag 436  
His Leu Thr Pro Phe Thr Ala Arg Gly Arg Phe Ala Thr Val Val Glu  
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Glu Leu Phe Arg Asp Gly Val Asn Trp Gly Arg Ile Val Ala Phe Phe

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Pro Leu Val Asp Asn Ile Ala Leu Trp Met Thr Glu Tyr Leu Asn Arg			
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gaa ctg tac ggc ccc agc atg cgg cct ctg ttt gat ttc tcc tgg ctg	676		
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Leu Gly Ala Tyr Leu Gly His Lys			
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Lys Tyr Ile His Tyr Lys Leu Ser Gln Arg Gly Tyr Glu Trp Asp Ala			
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Gly Asp Val Gly Ala Ala Pro Pro Gly Ala Ala Pro Ala Pro Gly Ile			
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Phe Ser Ser Gln Pro Gly His Thr Pro His Thr Ala Ala Ser Arg Asp			
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Pro Val Ala Arg Thr Ser Pro Leu Gln Thr Pro Ala Ala Pro Gly Ala			

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Ala Ala Gly Pro	Ala Leu Ser Pro Val	Pro Pro Val Val	His Leu Thr			
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Leu Arg Gln Ala	Gly Asp Asp Phe	Ser Arg Arg Tyr	Arg Arg Asp Phe			
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Ala Glu Met Ser	Arg Gln Leu His	Leu Thr Pro Phe	Thr Ala Arg Gly			
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Gly Arg Ile Val	Ala Phe Phe Glu	Phe Gly Gly Val	Met Cys Val Glu			
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Ser Val Asn Arg	Glu Met Ser Pro	Leu Val Asp Asn	Ile Ala Leu Trp			
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Met Thr Glu Tyr	Leu Asn Arg His	Leu His Thr Trp	Ile Gln Asp Asn			
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Gly Gly Trp Asp	Ala Phe Val Glu	Leu Tyr Gly Pro	Ser Met Arg Pro			
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Leu Phe Asp Phe	Ser Trp Leu Ser	Leu Lys Thr Leu	Leu Ser Leu Ala			
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 Leu Ser Tyr Lys Leu Ser Gln Lys Gly Tyr Ser Trp Ser Gln Phe Ser  
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 Asp Val Glu Glu Asn Arg Thr Glu Ala Pro Glu Gly Thr Glu Ser Glu  
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20080220 0630 020802

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gac agc ccc gcg gtg aat gga gcc act gcg cac agc agc agt ttg gat	362
Asp Ser Pro Ala Val Asn Gly Ala Thr Ala His Ser Ser Ser Leu Asp	
65 70 75	
gcc cgg gag gtg atc ccc atg gca gca gta aag caa gcg ctg agg gag	410
Ala Arg Glu Val Ile Pro Met Ala Ala Val Lys Gln Ala Leu Arg Glu	
80 85 90	
gca ggc gac gag ttt gaa ctg cgg tac cgg cgg gca ttc agt gac ctg	458
Ala Gly Asp Glu Phe Glu Leu Arg Tyr Arg Arg Ala Phe Ser Asp Leu	
95 100 105	
aca tcc cag ctc cac atc acc cca ggg aca gca tat cag agc ttt gaa	506
Thr Ser Gln Leu His Ile Thr Pro Gly Thr Ala Tyr Gln Ser Phe Glu	
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cag gta gtg aat gaa ctc ttc cgg gat ggg gta aac tgg ggt cgc att	554
Gln Val Val Asn Glu Leu Phe Arg Asp Gly Val Asn Trp Gly Arg Ile	
125 130 135 140	
gtg gcc ttt ttc tcc ttc ggc ggg gca ctg tgc gtg gaa agc gta gac	602
Val Ala Phe Phe Ser Phe Gly Gly Ala Leu Cys Val Glu Ser Val Asp	
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aag gag atg cag gta ttg gtg agt cgg atc gca gct tgg atg gcc act	650
Lys Glu Met Gln Val Leu Val Ser Arg Ile Ala Ala Trp Met Ala Thr	
160 165 170	
tac ctg aat gac cac cta gag cct tgg atc cag gag aac ggc ggc tgg	698
Tyr Leu Asn Asp His Leu Glu Pro Trp Ile Gln Glu Asn Gly Gly Trp	
175 180 185	
gat act ttt gtg gaa ctc tat ggg aac aat gca gca gcc gag agc cga	746
Asp Thr Phe Val Glu Leu Tyr Gly Asn Asn Ala Ala Ala Glu Ser Arg	
190 195 200	
aag ggc cag gaa cgc ttc aac cgc tgg ttc ctg acg ggc atg act gtg	794
Lys Gly Gln Glu Arg Phe Asn Arg Trp Phe Leu Thr Gly Met Thr Val	
205 210 215 220	
gcc ggc gtg gtt ctg ctg ggc tca ctc ttc agt cgg aaa tga	836
Ala Gly Val Val Leu Leu Gly Ser Leu Phe Ser Arg Lys	
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35 40 45

Ser Ala Ile Asn Gly Asn Pro Ser Trp His Leu Ala Asp Ser Pro Ala  
50 55 60

Val Asn Gly Ala Thr Ala His Ser Ser Ser Leu Asp Ala Arg Glu Val  
65 70 75 80

Ile Pro Met Ala Ala Val Lys Gln Ala Leu Arg Glu Ala Gly Asp Glu  
85 90 95

Phe Glu Leu Arg Tyr Arg Arg Ala Phe Ser Asp Leu Thr Ser Gln Leu  
100 105 110

His Ile Thr Pro Gly Thr Ala Tyr Gln Ser Phe Glu Gln Val Val Asn  
115 120 125

Glu Leu Phe Arg Asp Gly Val Asn Trp Gly Arg Ile Val Ala Phe Phe  
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Ser Phe Gly Gly Ala Leu Cys Val Glu Ser Val Asp Lys Glu Met Gln  
145 150 155 160

Val Leu Val Ser Arg Ile Ala Ala Trp Met Ala Thr Tyr Leu Asn Asp  
165 170 175

His Leu Glu Pro Trp Ile Gln Glu Asn Gly Gly Trp Asp Thr Phe Val  
180 185 190

Glu Leu Tyr Gly Asn Asn Ala Ala Ala Glu Ser Arg Lys Gly Gln Glu  
195 200 205

Arg Phe Asn Arg Trp Phe Leu Thr Gly Met Thr Val Ala Gly Val Val  
210 215 220

Leu Leu Gly Ser Leu Phe Ser Arg Lys  
225 230

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